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RAW SEQUENCE LISTING   DATE: 04/22/2002
PATENT APPLICATION: US/10/042,059A   TIME: 10:20:26

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF3\04222002\J042059A.raw**

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 3 <110> APPLICANT: Rhein Biotech Gesellschaft fur neue biotechnologische Prozesse und
 4        Produkte mbH
 6 <120> TITLE OF INVENTION: Nucleic Acid Molecule Containing a Nucleic Acid Coding for a
Polypeptide
 7        with Chorismante Mutase Activity
 9 <130> FILE REFERENCE: PA30558US-019
11 <140> CURRENT APPLICATION NUMBER: US 10/042,059A
12 <141> CURRENT FILING DATE: 2001-10-25
14 <150> PRIOR APPLICATION NUMBER: DE 199 19 124.7
15 <151> PRIOR FILING DATE: 1999-04-27
17 <160> NUMBER OF SEQ ID NOS: 3
19 <170> SOFTWARE: PatentIn Ver. 2.1
21 <210> SEQ ID NO: 1
22 <211> LENGTH: 843
23 <212> TYPE: DNA
24 <213> ORGANISM: Hansenula polymorpha
26 <400> SEQUENCE: 1
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28 cggatggagg atacgatcat cttcaacttt atcgagcggt cgcagttcta tgcgtcgccc 120
29 tcggtataca aagtcaacca gttccctatt cccaacttcg acggctcgtt cttggactgg 180
30 ctgttgtcgc agcacgagcg aatccattcg caggtgagga gatacgacgc gccagacgag 240
31 gtgccttttt tccccaacgt gctggaaaaa acgtttctgc ccaagatcaa ctacccatcg 300
32 gtgctagcct cctacgcgga tgaaatcaac gtcaacaaag agatactcaa gatctacacg 360
33 tcagagatag taccaggaat agctgcaggc agcggagagc aggaggacaa ccttggctcg 420
34 tgcgcaatgg ccgacatcga gtgcctgcag tcgctatcca gaagaatcca ttttggccgt 480
35 tttgtcgcag aggctaaatt tatcagtgag ggggacaaga ttgtggatct gatcaaaaag 540
36 agagatgtgg aaggcattga ggcgctcatc acaaacgccg aggtcgaaaa acggatcttg 600
37 gacagacttc tggagaaggg aagggcgtat ggaacagacc cgacactaaa gttcacgcag 660
38 cacattcaga gcaaggtgaa gcccgaggtg attgtgaaaa tctacaagga tttcgtgatt 720
39 ccgctcacga agaaggtcga agtcgactac ttgctgagac ggctggagga cgaggaggac 780
40 gatgatgcga cgcagaaaag cggcggctac gttgaccggt ttctctcctc tggcttgtac 840
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44 <211> LENGTH: 280
45 <212> TYPE: PRT
46 <213> ORGANISM: Hansenula polymorpha
48 <400> SEQUENCE: 2
49 Met Asp Phe Met Lys Pro Glu Thr Val Leu Asp Leu Gly Asn Ile Arg
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52 Asp Ala Leu Val Arg Met Glu Asp Thr Ile Ile Phe Asn Phe Ile Glu
53              20                  25                  30
55 Arg Ser Gln Phe Tyr Ala Ser Pro Ser Val Tyr Lys Val Asn Gln Phe
56          35                  40                  45
58 Pro Ile Pro Asn Phe Asp Gly Ser Phe Leu Asp Trp Leu Leu Ser Gln
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61 His Glu Arg Ile His Ser Gln Val Arg Arg Tyr Asp Ala Pro Asp Glu
62 65                  70                  75                  80
64 Val Pro Phe Phe Pro Asn Val Leu Glu Lys Thr Phe Leu Pro Lys Ile
65                  85                  90                  95
67 Asn Tyr Pro Ser Val Leu Ala Ser Tyr Ala Asp Glu Ile Asn Val Asn
68              100                 105                 110
70 Lys Glu Ile Leu Lys Ile Tyr Thr Ser Glu Ile Val Pro Gly Ile Ala
71          115                 120                 125
73 Ala Gly Ser Gly Glu Gln Glu Asp Asn Leu Gly Ser Cys Ala Met Ala
74      130                 135                 140
76 Asp Ile Glu Cys Leu Gln Ser Leu Ser Arg Arg Ile His Phe Gly Arg
77 145                 150                 155                 160
79 Phe Val Ala Glu Ala Lys Phe Ile Ser Glu Gly Asp Lys Ile Val Asp
80                  165                 170                 175
82 Leu Ile Lys Lys Arg Asp Val Glu Gly Ile Glu Ala Leu Ile Thr Asn
83          180                 185                 190
85 Ala Glu Val Glu Lys Arg Ile Leu Asp Arg Leu Leu Glu Lys Gly Arg
86          195                 200                 205
88 Ala Tyr Gly Thr Asp Pro Thr Leu Lys Phe Thr Gln His Ile Gln Ser
89      210                 215                 220
91 Lys Val Lys Pro Glu Val Ile Val Lys Ile Tyr Lys Asp Phe Val Ile
92 225                 230                 235                 240
94 Pro Leu Thr Lys Lys Val Glu Val Asp Tyr Leu Leu Arg Arg Leu Glu
95              245                 250                 255
97 Asp Glu Glu Asp Asp Asp Ala Thr Gln Lys Ser Gly Gly Tyr Val Asp
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100 Arg Phe Leu Ser Ser Gly Leu Tyr
101          275                 280
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104 <210> SEQ ID NO: 3
105 <211> LENGTH: 1655
106 <212> TYPE: DNA
107 <213> ORGANISM: Hansenula polymorpha
109 <220> FEATURE:
110 <221> NAME/KEY: gene
111 <222> LOCATION: (1)..(1655)
112 <223> OTHER INFORMATION: 1,8 kb genomic DNA-Fragment from Hansenula
113   polymorpha
115 <400> SEQUENCE: 3
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118 acgattacat tggcggaacg tatcgcgagt cgctcacgag acgcattaga atgacagaga 180
119 aatcgcgaaa cgaccttata gacgcacgtg aaaactacgg gtttggaggc agcaaggagg 240
120 agcgaatcca gcggtttttg tggttcagac atctttcgtg gcttttaggc gaggataagc 300
121 gaacttgagg agcgtttttt ttttcctgtt tagttttttgt aggtatggac tttatgaagc 360
122 cagaaacagt gctggacctt ggcaacatta gagatgcctt ggtccggatg gaggatacga 420
123 tcatcttcaa ctttatcgag cggtcgcagt tctatgcgtc gccctcggta tacaaagtca 480
124 accagttccc tattcccaac ttcgacggct cgttcttgga ctggctgttg tcgcagcacg 540
125 agcgaatcca ttcgcaggtg aggagatacg acgcgccaga cgaggtgcct tttttcccca 600
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126 acgtgctgga aaaaacgttt ctgcccaaga tcaactaccc atcggtgcta gcctcctacg 660
127 cggatgaaat caacgtcaac aaagagatac tcaagatcta cacgtcagag atagtaccag 720
128 gaatagctgc aggcagcgga gagcaggagg acaaccttgg ctcgtgcgca atggccgaca 780
129 tcgagtgcct gcagtcgcta tccagaagaa tccattttgg ccgttttgtc gcagaggcta 840
130 aatttatcag tgaggggac aagattgtgg atctgatcaa aaagagagat gtggaaggca 900
131 ttgaggcgct catcacaaac gccgaggtcg aaaaacggat cttggacaga cttctggaga 960
132 agggaagggc gtatggaaca gacccgacac taaagttcac gcagcacatt cagagcaagg 1020
133 tgaagcccga ggtgattgtg aaaatctaca aggatttcgt gattccgctc acgaagaagg 1080
134 tcgaagtcga ctacttgctg agacggctgg aggacgagga ggacgatgat gcgacgcaga 1140
135 aaagcggcgg ctacgttgac cggtttctct cctctggctt gtactagaaa ttaaaatttt 1200
136 cagtacttta attattctcg aattctagtt cagataccgc atggtaattt caaaggccag 1260
137 aaaagtggcc gcgttggctg gggcagctct cagaatagtc ggcgagaatc ctttgactag 1320
138 cccccaggca ccgctctgtc tccaaatacc cctaatagtc tcaacagcat ttctataaac 1380
139 cagcttcttg tagttgtccg tctgcatgtt ggacttgatc acatcgatcg gataaatact 1440
140 gaaccacatc ccgtaacctg ccagcgcccc aaagacgcag agcttccagt tctcgatgtc 1500
141 cttcctggca atattccgcg actcgatctc gtttttcacg agagcttcaa aagtcagaaa 1560
142 atacgctccg ctacccaaac tttctcttgc cagcgtaggt cccagacccc ggtagattaa 1620
143 cttgatgcct cccgtatggt acagcttctt gatcc                             1655
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